**S2 Table: Mean absolute error (MAE) segmentation regression estimates following cross-validation along with use of lesion masks (mean sd).**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Tissue** | **Blue (Myelin)** | | **Red (Cellularity)** | |
| **Intercept** | **Slope** | **Intercept** | **Slope** |
| SG |  |  |  |  |
| GM |  |  |  |  |
| PV |  |  |  |  |
| WM |  |  |  | 0 |
| Lesion |  |  |  |  |